

SEQUENCE LISTING

<110> Kenneth W. Dobie

Mark P. Roach

<120> ANTISENSE MODULATION OF NOD1 EXPRESSION

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aact atg gaa gag cag ggc cac agt gag atg gaa ata atc cca tca gag     469
      Met Glu Glu Gln Gly His Ser Glu Met Glu Ile Ile Pro Ser Glu
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Ser His Pro His Ile Gln Leu Leu Lys Ser Asn Arg Glu Leu Leu Val
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act cac atc cgc aat act cag tgt ctg gtg gac aac ttg ctg aag aat      565
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cag cct gac aag gtc cgc aaa att ctg gac ctg gta cag agc aag ggc      661
Gln Pro Asp Lys Val Arg Lys Ile Leu Asp Leu Val Gln Ser Lys Gly
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gag gag gtg tcc gag ttc ttc ctc tac ttg ctc cag caa ctc gca gat      709
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Ala Tyr Val Asp Leu Arg Pro Trp Leu Leu Glu Ile Gly Phe Ser Pro
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Arg Tyr Thr Gln Gln Leu Arg His His Leu Gly Arg Asp Ser Lys Phe	
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Val Leu Cys Tyr Ala Gln Lys Glu Glu Leu Leu Leu Glu Glu Ile Tyr	
145 150 155	
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Met Asp Thr Ile Met Glu Leu Val Gly Phe Ser Asn Glu Ser Leu Gly	
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Ser Leu Asn Ser Leu Ala Cys Leu Leu Asp His Thr Thr Gly Ile Leu	
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Asn Glu Gln Gly Glu Thr Ile Phe Ile Leu Gly Asp Ala Gly Val Gly	
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Lys Ser Met Leu Leu Gln Arg Leu Gln Ser Leu Trp Ala Thr Gly Arg	
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Leu Asp Ala Gly Val Lys Phe Phe Phe His Phe Arg Cys Arg Met Phe	
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Lys His Tyr Cys Tyr Pro Glu Arg Asp Pro Glu Glu Val Phe Ala Phe	
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Leu Leu Arg Phe Pro His Val Ala Leu Phe Thr Phe Asp Gly Leu Asp	
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Glu Leu His Ser Asp Leu Asp Leu Ser Arg Val Pro Asp Ser Ser Cys	
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ccc tgg gag cct gcc cac ccc ctg gtc ttg ctg gcc aac ctg ctc agt	1381
Pro Trp Glu Pro Ala His Pro Leu Val Leu Leu Ala Asn Leu Leu Ser	
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Gly Lys Leu Leu Lys Gly Ala Ser Lys Leu Leu Thr Ala Arg Thr Gly	
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Ile Glu Val Pro Arg Gln Phe Leu Arg Lys Lys Val Leu Leu Arg Gly	
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Phe Ser Pro Ser His Leu Arg Ala Tyr Ala Arg Arg Met Phe Pro Glu	
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Arg Ala Leu Gln Asp Arg Leu Leu Ser Gln Leu Glu Ala Asn Pro Asn	
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Cys Phe Gln His Phe Arg Ala Ala Phe Glu Gly Ser Pro Gln Leu Pro	
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Asp Cys Thr Met Thr Leu Thr Asp Val Phe Leu Leu Val Thr Glu Val	
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His Leu Asn Arg Met Gln Pro Ser Ser Leu Val Gln Arg Asn Thr Arg	
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Ser Pro Val Glu Thr Leu His Ala Gly Arg Asp Thr Leu Cys Ser Leu	
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 Gly Gln Val Ala His Arg Gly Met Glu Lys Ser Leu Phe Val Phe Thr
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 Gln Glu Glu Val Gln Ala Ser Gly Leu Gln Glu Arg Asp Met Gln Leu
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 Gly Phe Leu Arg Ala Leu Pro Glu Leu Gly Pro Gly Gly Asp Gln Gln
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 Ser Tyr Glu Phe Phe His Leu Thr Leu Gln Ala Phe Phe Thr Ala Phe
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 Pro Pro Phe Leu Pro Phe Gln Cys Leu Gln Gly Ser Gly Pro Ala Arg
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 Glu Asp Leu Phe Lys Asn Lys Asp His Phe Gln Phe Thr Asn Leu Phe
 580 585 590

ctg tgc ggg ctg ttg tcc aaa gcc aaa cag aaa ctc ctg cgg cat ctg 2245
 Leu Cys Gly Leu Leu Ser Lys Ala Lys Gln Lys Leu Leu Arg His Leu
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 Val Pro Ala Ala Ala Leu Arg Arg Lys Arg Lys Ala Leu Trp Ala His
 610 615 620

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 625 630 635

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Ala Arg Gly Ile Cys Ala Asn Tyr Leu Lys Leu Thr Tyr Cys Asn Ala	
675 680 685	
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Ile Thr Asp Val Gly Ala Arg Tyr Val Thr Lys Ile Leu Asp Glu Cys	
770 775 780	
aaa ggc ctc acg cat ctt aaa ctg gga aaa aac aaa ata aca agt gaa	2821
Lys Gly Leu Thr His Leu Lys Leu Gly Lys Asn Lys Ile Thr Ser Glu	
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Gly Gly Lys Tyr Leu Ala Leu Ala Val Lys Asn Ser Lys Ser Ile Ser	
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 Phe Ala Glu Ala Leu Arg Asn His Pro Ser Leu Thr Thr Leu Ser Leu
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 Gln Thr Leu Lys His Leu Trp Leu Ile Gln Asn Gln Ile Thr Ala Lys
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 915 920 925

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 Tyr Glu Asp Glu Lys Arg Ile Ile Cys Phe
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<212> DNA

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1. The invention relates to a method for determining the sequence of a nucleic acid molecule, comprising the steps of: (a) providing a nucleic acid molecule; (b) sequencing the nucleic acid molecule; and (c) determining the sequence of the nucleic acid molecule.

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Ser Gly His Ser His Pro Gln Tyr Ser Val Ser Gly Gly Gln Leu Ala
15 20 25 30

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Glu Glu * Leu Leu Leu Gly Gly Arg Cys Gly Asp Cys Val Cys Leu
35 40 45

ccc cac cca gcc tga ca ggtgccccgg ggacaggac gggcatgggc 191
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